

SEQ ID NO:1

1 GGCAATGGGT CGAAATTCAT AGAATTTTGT GTGAGGTGCG TAGCGGCTCT  
51 GACAGGGGTG CTGCGCGGAG ATCTCTGGTC TCAGGTAGGG CGACAATGGA  
101 GAGGTGTTAG TTGCCCCCTG TATCGCTCTC TCGGTGGCGC ATTGGGTCAT  
151 CCTGCCCCGA CATATGATAT TCCGCTAGAG GATTACTGAT AGTTTCTGCC  
201 TGTCGGGCTT GTCGGGCTTG TCGGGCTTGT CGGGCTTGTC GGGCCTGTCC  
251 CTCTTGTCCT GCCTGTCCTC ACTTTTTCAC AATCAAAAAA TGGGCGAAGC  
301 CCTTCTTGTT CTATAGTTCT TATAGTTCAT ACGAAAATTA CACATAATTA  
351 TCAATAGCTT ATTCGCTTAA AAGGGAGTAA TTGGGCCGCA AAAGGGAGTA  
401 ATTGGGCCGC AAAAGGGAGT AATTGGGCCG CAAAAGGGAG TAATTGGGCC  
451 GATATCGGTT GTTTACATGG GGAGGAATCC CCTTAATCAT TTCTCCCCAT  
501 GGGAAAGACA ACACAAGTGG CCGCAGACCG GGCCTTCGAC CAGACAAAAA  
551 CTGTGCTCCC TGCCGAGGTG GCGAGAGGGG TCTATATGCG CAATCCGCCC  
601 CGCCTGCAGG CGCTCAAGCT CATGCATTTA ATGATAGCCA CTGCGGGCGG  
651 CCGCATGGCT GATGATGTGC GCCATGAAAT GCGGCTGGCC GACATTGCGG  
701 CAATCGACGG CATGAAAAAC CATGACCGTG AGAGCCTGAC CCCGCTGTTC  
751 GAGGAGCTAG CCGCTGCGGT GTTGACCCAT GATGACCTG CAAAGATGAT  
801 CGTGACAGTC GGCGGCTTGG TCGATGAGGC GCGAATAGAC TACCGCCAGG  
851 AGGCAAGCGG CGAACTCCTA GTGACGTGGA CCTTCCGGAG TACATTCCGT  
901 CGTATGGCGG CGGAGTCGAA CCACTGGGCC ATTCTCGACC GTCAAACGGT  
951 ATTCCATCTC GGTAGTAAGT ATTCCGTGCT GCTGTTCCAG CACGTCTCTA  
1001 GTCTCGCCAA TCTTGATCGG ATGAGCGCGA AAACCTTTAC GGTCCCCGAG  
1051 TTGCGGGCGC TCCTTGAGT GCCCGAGGGA AAGATGGTTC GTTGGAACGA  
1101 CGTTAACAGA TTTGCTCTCA AACCTGCACT GGATGAGATC AACCATTAT  
1151 CGCGTCTGAC ATTGACGGCA AAGCCGACCA AGATTGGCCG TAGCGTGGA  
1201 AGTGTGACTA TAGGCTGGGA AGTGAAAGAC GACCCAACCG TCGCCAGGCG  
1251 CGAGCTGGCG GGTTCOAAG TCGGTCGAGA TGCTCGTCGC AGAGGGGCAG  
1301 CGGAAACGAT AGCCCCCTCC TTCCAGAAAG CGGGCGGGAT CACCTACAGT  
1351 CCACGTTGGC TGGAGCTGAA ACGCTCTGCT GGCAGCAACA AGGACAACGA  
1401 TCTGATCGCC TCAGACTTCC GCGGTTTCTG TCGGGAGAGA GCGGTGCGTC  
1451 TGGACGCTGC AAACATCGAA AAAGTGTGTT TAGATTTCTG CGCAAAGGTA  
1501 GGAAGGTTT GAGTTTGTAG GTATTTTACC GCAATAGTGT TAAATGACTT  
1551 TCGTGAAACG ATGTGCAATA TAGCGGTAAG ACTATGAAAT ACACGGCTGG  
1601 ACAGGCTGCA AAAGCAACGG GTGTGGCGAC CGCAACCATC ACTCGGGCGC  
1651 TAAAAAGCGG TAAAATTTC GGTAAAAAAG ATGAATCTGG GGCATGGGTT  
1701 ATAGATCCTG CAGAAATGCA CAGAGTGTTT CCTCCCATT CAAAGAAATA  
1751 CACCGAAACA CTAACACGC AAGTATATGG TAAGCGTGAT GAAACACATG  
1801 AAATGACCTC AGAAATCAGC GCATTAGAGC GTGAAGTTCG GACTTTACGC  
1851 GATGCTTTAT CTGATGCCAG GGAGGATCGC GACAAATGGC GCGACATGGC  
1901 CGAGCGTCTT TCAATTTTAT CACCGATGAG AGAGGAAGAC CGCCCCCTC  
1951 AAAACAAAG ATGGTGGAAG ATATTCTGAT CCTGGGCTTC AGGAGCCTTG

FIGURE 1A

[illegible]

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SEQ ID NO:2

1 TGGTGAACGC ATTGGCTTGA TGTTTGAGAA AAGCGAAAAG ACCCGGCCAC  
51 AGTTGTGGGT AGAGCGTCGA TATGTGCAAG ACCTGATGCT TGCTGACATC  
101 GAACTCCGTG TCTACCTCGC ATCGTCGCTG TATCAGCCTG CTGCGGATGG  
151 CGGAAAGCCC GCCTATGGTC GTCACGCAGC CCTTAAGGCG ATGCGCGACT  
201 TGGCCCATGC CGATCTGGTG CGTTTCACCA TCGGCCGGAT TACGCAACTG  
251 GAGATGATCC TAGAGCGGTT AACCAGAGACA TCTGGTTAAC GCCATAAAGG  
301 CTGCGGCATG AAAATAGGCG GACAATCTGC GCTTGGCCGC CCCCGTTCTC  
351 AGCCGTGCTT GCTCTCTGCC TGCATGGCAC GACGCAGGAT CGCGTTTATA  
401 CGGGTCTGAT ATCCAGACCC GCCCGCCTTG AGCCATGCCA GCACATCGGC  
451 ATCAAGCCGC GCGGTGATCT GCTGCTTGAT CGGGCGATAG AAGCGCCAC  
501 GCTCGGCGTC TGCCCATTTG GCTTCGGTCA GCTCGGGAAC ATCGTTGGTG  
551 TCGATCTGCT CGGGCGGCAG AGCGTCCAGC CGCGCCAATT TCTTGCGGCG  
601 CTCTCTGGTA AGAGCGGGCA GCGTATCGAA GGTGTATTCA ACCATTGGCA  
651 TATCTCTTCC TTTCTGCGG TGTAGCGCGG CGAGCCGAAA TGATGCGGAT  
701 CGTCTCGACC GGATCGGGGC CAGCCTCGAT GATCAGGTGG GCAACCAGAA  
751 GGACGGCAGC GCCATAGATC TGCCCAACGG TTTGCCAGCG GTATTCCCCG  
801 CCCTCGATCC TATCTGAAC CGTCAGGTGC AACGGATCGG CGAACACATG  
851 CACAGCATCC TCGAACCGBA TGCCATGCTT CTTTTCGTTT GTTTCCGCTT  
901 TGGCGGGATC CCAGATAAAC CGCATCTTCA TGGCAGAATT ATAACCTACAC  
951 ATTTGTAGTT ATTCAATGGC AAGTCGCAGG TTCAAATCAC GCCCCCAAAC  
1001 CGCAACTGTA TTCGTTCTAC TCACGCGCGC TTTTGAATAG AAGCTTGCAT  
1051 GATAACACCC GCCGCGTCCT CAACAAAATA AGGCAAATCC GCCGCGCTGG  
1101 CGCAATCTGC GCTTTGTCGA TGCAAGGTCT TGTGGTTTCA TACTGCAAGA  
1151 GCATGCAAGG AATTGCCCCG GATGAGCACC ACGACGACAC CCACCAAGCC  
1201 GGCTTGAAC AAGGGCCGCG TTGTGCGGAA AAAGCCGCCG CTGACACCTG  
1251 ACCAGATTGC CCTGATCCGT CTCATCCTGC GCCAGGAACG GGCCTGGCGG  
1301 GATCTGGCTC GTTCAACGT GCGATCGAC ACCAGTTTGC GCGGCTCGGA  
1351 CCTCGTGCGC CTGCGCGTCT CGGATGTGGC GACCCAGCT GGTCTGCGTG  
1401 AGATCGTCGA GATCCGCCAG AAGAAGACCG AGGCCCGCAA TGTCCGCCCC  
1451 GTACAGGCCC GCCTGTCGGA GGGGACACGC GAGAGCCTGC GGGTCTATCT  
1501 CGCGGCCTCT GACAAGCCGC TGCACAGCTG GCTGTTTACC GGACAGGGCA  
1551 TCCGCTGGTC CCACACCCAC CTTAGCGAGA GCCAGCTGTG GCGCCTGTTT  
1601 AAGTCCTGGC TCGAGAAGGC GCGGCTCGAT CCCAGCCTCT ACGGGCTGCA  
1651 CTCGCTGCGC CGAACCTTCC CCAGCCACAT CTACCGCGAG ACCGGCAATC  
1701 TGCGCGCCGC ACAGCTGCTG CTGGGCCATG CCAGCATCGA GAGCACCAG  
1751 GAGTACATCG GCACCGAGCA AGCCGAGGCC CTCGATATCG CACGGAGGTA  
1801 TCACCTCTAA CCCATGGAGA CCTATCTCGA GAAGCGCATC CCCGCCAAGA  
1851 ACACAGCACG GTTCTACCGC ATGGCGGTCC TGCCGAACCT GTTCGGGGAA  
1901 TGGACGCTGT ATCGAGAATG GGGCCGCATT GGCATCAGCG GCCGCATCCG  
1951 GCTCGATTGG TTTGAGAGTG AACAAGATGC CATCGCTGCG ATGCTCGCCA

FIGURE 2A

2001 TCGAGACCGC CAAGCGTCAG CGCGGGTATT GGCTCGAGCC CATCCAGATT  
 2051 GACATGTTCC CAGGGGCATA ACAGGCCATC AATGTAAGAG TGCAAGCGGA  
 2101 GCAAGCAAAA GCCATTTTAC AGTGAGGTGG CAGATGTTCC TGTTCACAG  
 2151 TGAAAGCGCT GATGCTGTTT CCACGCCACA GACTGATACG ACCAAAGCAA  
 2201 CGGGGTCTGC CGCCACAGAC CGGTTCGCCG GCCACCCGCA GAAACGCAGG  
 2251 TAAAATGGCG ATTTCCGCAA AAAAACCGTG CAAATGATGG CAAATCACCA  
 2301 TCCAGTTTCA TCCTGAAACC CGTCGCTCAA CATGAACGAG CAGGCCATCA  
 2351 TCCAAGCCCC AGAAACGCGG TGCGGCGACT ACAGATGAGC GATGTTCTGG  
 2401 CTCATAGGCT GCAAGGCCCT GCAACAGTGA TTTCACCGTG AGATTGCAGG  
 2451 GTCTTTTGGC TCTCCCGCAA GAGCCACCTC AGGGTGAGCG AGCTAGCCGT  
 2501 CTAGGTTTAC AGTGAAATCG CTGAGGAGCG TTGCGGGGCT TATGGTTTGG  
 2551 CTGGTCACGT TGGCCATCGG AATGGAGCAT ACGATGGCTT CTACGCAGTC  
 2601 GAATCCTGAG GCTTCACGTG GGAAAAATAC GCTCCAAAAA AGCCCTGACC  
 2651 AAATCTTGGA AAAATTGCTT GAAAAGTTTG CTTCTAAAAA ACTGGGAACG  
 2701 AGATATGCAC GAGATCCCTT ACGAGTGCTG TAGGAGTAAT GCAGTGGACA  
 2751 AAAACGCCAT TTTTGTCCCC AGTAGGAGTA ATGGAGTGGT TATTTTTTGG  
 2801 GAGATTTTGC TTCAGTAGGA GTAACGCGTT GGTAAATTT GCTTGATTGG  
 2851 CGGTTCAAAT CGACCACCGA GCTGCCGTTG GTCGTATTCG ATCTGCCCCG  
 2901 CAATTGGGCA CTTGCAGGCC ATCCCCCTGA ACTTCTGGCG ATGACCATTT  
 2951 CGAAGGCAAT GGGTCGAAAT TCATAGAATT TTGTGTGAGG TGCGTAGCGG  
 3001 CTCTGACAGG GGTGCTGCGC GGAGATCTCT GGTCTCAGGT AGGGCGACAA  
 3051 TGGAGAGGTG TTAGTTGCCC CCTGTATCGC TCTCTGCGTG GCGCATTGGG  
 3101 TCATCTTGCC CGGACATATG ATATTCCGCT AGAGGATTAC TGATAGTTTC  
 3151 TGCCTGTCGG GCTTGTGCGG CTTGTGCGGC TTGTGCGGCT TGTCGGGCGT  
 3201 GTCCCTCTTG TCCCGCCTGT CCTCACTTTT TCACAATCAA AAAATGGGCG  
 3251 AAGCCCTTCT TGTTCTATAG TTCTTATAGT TCATACGAAA ATTACACATA  
 3301 ATTATCAATA GCTTATTCGC TTAAAAGGGA GTAATTGGGC CGCAAAAGGG  
 3351 AGTAATTGGG CCGCAAAAGG GAGTAATTGG GCCGCAAAAG GGAGTAATTG  
 3401 GGCCGATATC GGTTGTTTAC ATGGGGAGGA ATCCCTTAA TCATTTCTCC  
 3451 CCATGGGAAA GACAACACAA GTGGCCGCAG ACCGGGCCTT CGACCAGACA  
 3501 AAAACTGTGC TCCCTGCCGA GGTGGCGAGA GGGGTCTATA TGCGCAATCC  
 3551 GCCCCGCCTG CAGGCGCTCA AGCTCATGCA TTTAATGATA GCCACTGCGG  
 3601 GCGGCCGCAT GGCTGATGAT GTGCGCCATG AAATGCGGCT GGCCGACATT  
 3651 CGCGCAATCG ACGGCATGAA AAACCATGAC CGTGAGAGCC TGACCCCGCT  
 3701 GTTCGAGGAG CTAGCCGCTG CGGTGTTGAC CCATGATGAC CCTGCAAAGA  
 3751 TGATCGTGAC AGTCGGCGGC TTGGTCGATG AGGCGCGAAT AGACTACCGC  
 3801 CAGGAGGCAA GCGGCGAACT CCTAGTGACG TGGACCTTCC GGAGTACATT  
 3851 CCGTCGTATG GCGGCGGAGT CGAACCCTG GGCCATTCTC GACCGTCAAA  
 3901 CGGTATTCCA TCTCGGTAGT AAGTATTCCG TGCTGCTGTT CCAGCACGTC  
 3951 TCTAGTCTCG CCAATCTTGA TCGGATGAGC GCGAAAACCT TTACGGTCCC  
 4001 CGAGTTGCGG GCGCTCCTTG GAGTGCCCGA GGGAAAGATG GTTCGTTGGA  
 4051 ACGACGTTAA CAGATTTGCT CTCAAACCTG CACTGGATGA GATCAACCAT

**FIGURE 2B**

4101 TTATCGCGTC TGACATTGAC GGCAAAGCCG ACCAAGATTG GCCGTAGCGT  
4151 GGCAAGTGTG ACTATAGGCT GGAAGTGAA AGACGACCCA ACCGTCGCCA  
4201 GCGCGAGCT GCGGGTTCC AAGGTCGGTC GAGATGCTCG TCGCAGAGGG  
4251 GCAGCGGAAA CGATAGCCCC CTCCTTCCCA GAAGCGGGCG GGATCACCTA  
4301 CAGTCCACGT TGGCTGGAGC TGAAACGCTC TGCTGGCAGC AACAAAGACA  
4351 ACGATCTGAT CGCCTCAGAC TTCCGGCGTT TCTGTCGGGA GAGAGGCGTG  
4401 CGTCTGGACG CTGCAAACAT CGAAAACTG TTTTATAGATT TCTGCGCAAA  
4451 GGTAGGGAAG GTTTGAGTTT TGAGGTATTT CACCGCAATA GTGTTAAATG  
4501 ACTTTCGTGA AACGATGTGC AATATAGCGG TAAGACTATG AAATACACGG  
4551 CTGGACAGGC TGCAAAAGCA ACGGGTGTGG CGACCGCAAC CATCACTCGG  
4601 GCGCTAAAAA GCGGTAAAAA TTCCGGTAAA AAAGATGAAT CTGGGGCATG  
4651 GGTATATAGAT CCTGCAGAAT TGCACAGAGT GTTTCCTCCC ATTTCAAAGA  
4701 AATACACCGA AACACCTAAC ACGCAAGTAT ATGGTAAGCG TGATGAAACA  
4751 CATGAAATGA CCTCAGAAAT CAGCGCATTG GAGCGTGAAG TTCGGACTTT  
4801 ACGCGATGCT TTATCTGATG CCAGGGAGGA TCGCGACAAA TGGCGCGACA  
4851 TGGCCGAGCG TCTTTCAATT TCATCACCGA TGAGAGAGGA AGACCGCCCC  
4901 CCTCAAAAAC AAAGATGGTG GAAGATATTC TGATCCTGGG CTTCAGGAGC  
4951 CTTGCCTTTA AAACCTGAAT CAGCATTCTA GCGATGCTGA TAAGAAGTAA  
5001 ATATAGCCAC AATAGAGCGG CCATTTTCCA TTCACATACA GCTCATCATG  
5051 TGATCAATAT CAAGTATTGA TATTCATCAA TGGAGAAGAA TTTACATGTA  
5101 TCACAGGATC ATCACAGCAT TTGTTTTTGT ATTTCTAAGT GCTAACATAA  
5151 CTATCGCTGG CCTTAAAGAA GATTGTACTA TTGCAGTATC TCACCTTGGG  
5201 TTTCAGACCG ATAATTACAG CTTTGTGCGA GCCGGTTTTT TTGCCAGAGA  
5251 GAGACACGTT TTTGATGGTG TAATAAACTG CTACGTATCT CATGATGGTA  
5301 ACATACACAG CATCATCCGG GGCAACACAC CTCTTATGGA AGATGGATAT  
5351 TATGGCCCAG AAGTACTGGC GGAAAAACGC GATATTGAGG CACAGGCCCC  
5401 CACTTTAGAG GCGGAAGCCT ATAACGAGTA CCAAAACACT AGAAGCCAGA  
5451 TTGAGGAAAA TAGGGAACGT GCCCTCGAGG CGCTGCGGCT AGCTAGCAGT  
5501 CCTTTTATTA ATAATGGTAG TACAGAAGAA CAGACAATTA TACAGGCCGC  
5551 AACTCCGACG GCAGATCCTG TTGTATCTGT ACCCGTGGCA TCTCCAGAAT  
5601 CTAAACAAAG TCGAGAGCCG GAACCGGCTG CTGTTCCAGC ATCAGTTTCT  
5651 GTTAGAGAGA TGTGGAGCAC GGCTGACAGA TTGACCACCC GTACATGCCC  
5701 ATCGACTCGA TGCGGAGCAA CTAGCTGGGT AACAGATGGA ACTAAAGTAA  
5751 CAGTTTATGA AGAAAAAGAC GGTGGGTCTA GAATCGGAGA GCTACAGTCT  
5801 GCAATGTGCA TAAATGGAAT AAGTGGCGCG GTCGATTAG GTGAATCTTC  
5851 CTGCAATCCC ACCAATGGTA TCGTTAATGG GCAATTCGCA CCCTGGGTTT  
5901 TCTCGGATTA TCTTACGATC CAAGAGCCAG AAGCTCCCAT ATCCACCCAA  
5951 GAGTGTGCGA ATATGGGGCT CGAGAACTCA GATAATTACC GTATCTATTC  
6001 TAGTCAGTTC TGCACTGCCG CTCTCGAAAT GATCAACGAT AGAGTATGCA  
6051 ATACATCTGA TTTCAGAGAT TTAGCTTGGT TATCTTCTCC TGAAAGAGGA  
6101 CAGGATTACT ACTTCACCTA TTGTGGCGGA TTTCAACCTC AAAACAGATG  
6151 GTATTTGAAT GTCAGGACAG GTGAAATCAC CCGCTGATAT TCCACCAAGG

**FIGURE 2C**

6201 TGAGTCCTGT AGATCAGACT CTCAAGGAGT AAACGTTTTTA ATCCATCTCC  
 6251 ATGAGATCAA CATAGATAGG TGTTCAGTCC CGGCATCTGG TGGATCGGGT  
 6301 TTAGGATGAA TCTGTCCGGC TCTTGACATA CCCCCGCGTG AAACCCGTGC  
 6351 TTTACAAGAG AAAGTCAGCG GCCTCGAAGC CGCTCTAGCC GATGCCCCGG  
 6401 CCCAACGGGA TGAGTAGAGC GAACAAGCAA AGCGCCTAGC TATGGCTCTG  
 6451 CCCGTCCCCG AAGCTGCAGC CGCAGAATCC GGAAAAAGA AAAAATACAT  
 6501 GGCAGCGATT ATTTGGATAG GACACAATCC TTTTCTATTA ATATACAACA  
 6551 AGATATGGGC ATGCGCCGCG CGTGATCCTC ATTGCATACA ATCCAAATCC  
 6601 TGAAAGCTGA CTATGCCCTA CGCATCGCGC ACCATCGGTG CCGTCATTGA  
 6651 TGACGTGAAC CGCACCTACC TGCTGCCCCG AATCCAACGC CCCATGTCTT  
 6701 GGTCTGCCGG ACAGGTCGTT GCGCTGTTCT ACTCTCTGTT GAAGGGCTAT  
 6751 CCGATCAGCA GCTTCATGTT CTGGGCGGTG GACGAGGAGA CCAAGGCAGA  
 6801 GCTGCGATGC TACAAATTCA TCGAGAATTA TCGGCCCGAA ATGATGAACG  
 6851 AGCCGACTAG TGCGGACGGG CGGCAGGTCG TCCTTGTGCT CGACGGACAG  
 6901 CAGCGGATGA CCTCACTGTT GATCGGCTTG CGCGGCACAT TCTCTGAGAA  
 6951 AGCCAAACAC GCGCGCAACA GCAACGCGGC GGCGTGGTCG GCAAAAACGC  
 7001 TATATCTAGA CCTGCTTCGG GACCCGGATC CGAAGAACTC CGATGAAGAC  
 7051 GAAGGCAATG AGTTCGGAAT CACTTACGGT CTCTCTTTCC ATGAACGCCG  
 7101 CCCGACCAGC AGCCACAGGC ACCACTGGTT CAAGGTGGGA TCGATACTGG  
 7151 ATTATCCTAC AGACGAGCAG CTGGAGGGGT TGATTGCCAA GGTGAAGACC  
 7201 GAATTTTCATC ATGGTGTATC GGATTGGGAA AAGGGGCTGG CGGAAGACAC  
 7251 CCTGCGCCGG TTGCACCGCG TCATCTGGAA AGACGAGGGC ATCAACTTTT  
 7301 TCACTGAACG CGACCAGTCG GTTGATCGGG TGCTGGACAT CTTCGTGCGG  
 7351 GCCAATGACG GGGGCACGAA ACTGTCGAAG GCAGACCTGC TGATGTGCAT  
 7401 GATCACGTCA AAATGGTCCA GCGGATCGGC CCGCGAGGAA ATCGGCGGCT  
 7451 TTGTCGAGCA CATAAACAAA GGTCTCGGTG CGCCGAACAA GATCAGTCGC  
 7501 GATCTGGTCC TGAAGGCCTG TCTGGTCGTC TGCGATTATG ATGTCGTCTA  
 7551 TAATGTCAGG AACTTTACAA GCGAGGTCAT CGGCAGGATC GAAAGCAACT  
 7601 GGGATCGTAT CAAGCAGGCA TTCGAGAACA CGTTCCGCCT GCTGAACAGG  
 7651 CATGGCATCA CCGGGGATAA CCTCGGCTCT TTGAACCGCG TGCTGCCTCT  
 7701 GGTCTATTAT ATCTACAACA CGCCGGATTT CGATTTCCGA GGATCGAGCG  
 7751 AGTTCGAGCG GGTCAATGCC AGCTCCATGC ACCTCTGGTT GGTGAACAGC  
 7801 CTGCTGGTCA GCGCCTTCGT TGGCCATTCG GATCAGACCA TCACCACCGC  
 7851 GCGCAATACG ATCCGCGATC ACCTGCGTGT AGGCCGCGAT TTCCCAGTAC  
 7901 GAAAGCTGTT CGATGCCATG GCGAAGGGGG GACGGCTATC TCAGGTGGAT  
 7951 GAGCGTACCA TCGAAGAATT GCTGGAAATG CAATATGGCA AGCCCCGGAC  
 8001 CTTCGTTGCG CTGTCGCTGC TCTATCAGGG CATCGACTGG AACGGATCGA  
 8051 CCTGGCATGT CGATCATATC ATTCCCCAAG CGGACGCTCA GAAAAATGTG  
 8101 CTGCGCGGGC GCAATCTGCC CGAGCATCGC ATTCAGGAAA TCTTGGGCGC  
 8151 GGTAAACAGT TTGGGCAACC TGCAACTTTT GCGCGGAGAT GAGAATATCG  
 8201 AGAAAGGTGC GCTGCCATTC AGGTCATGGA TTACCGGACG GCGCGTTGAT  
 8251 TTCTACGAGC AGCATATGAT CCCGCGCAC CTTGAACTGT GCGATGTACT

**FIGURE 2D**

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8301 GCATCTGCCC GAGTTCGTGC GCGAACGGGA AAAGGTGATC CGGCGCCGTT
8351 TGATGGAGTT GGTCCGAGCA CGACGCGCAT GAATGAGGTC GTCTTGTCAC
8401 GCGAAGAGCT GCGTCAATCT TGTCTCGACC TGCTTGAAAA ACGCGCTGTC
8451 GAACACCCTG CGGGACACCA AGGCAAGCTC GCCGCCCCGCT ATGTTGTGCA
8501 CCGCGACGA

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**FIGURE 2E**

8301 GCATCTGCCC GAGTTCGTGC GCGAACGGGA AAAGGTGATC CGGCGCCGTT  
 8351 TGATGGAGTT GGTCCGAGCA CGACGCGCAT GAATGAGGTC GTCTTGTCAC  
 8401 GCGAAGAGCT GCGTCAATCT TGTCTCGACC TGCTTGAAAA ACGCGCTGTC  
 8451 GAACACCCTG CGGGACACCA AGGCAAGCTC GCCGCCCCGCT ATGTTGTGCA  
 8501 CCGCGACGA

SEQ ID NO:3

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1  TCGCGCGTTT CGGTGATGAC GGTGAAAACC TCTGACACAT GCAGCTCCCCG
51  GAGACGGTCA CAGCTTGTCT GTAAGCGGAT GCCGGGAGCA GACAAGCCCCG
101 TCAGGGCGCG TCAGCGGGTG TTGGCGGGTG TCGGGGCTGG CTTAACTATG
151 CGGCATCAGA GCAGATTGTA CTGAGAGTGC ACCATATGCG GTGTGAAATA
201 CCGCACAGAT GCGTAAGGAG AAAATACCGC ATCAGGCGCC ATTCGCCATT
251 CAGGCTGCGC AACTGTTGGG AAGGGCGATC GGTGCGGGCC TCTTCGCTAT
301 TACGCCAGCT GCGGAAAGGG GGATGTGCTG CAAGGCGATT AAGTTGGGTA
351 ACGCCAGGGT TTTCCAGTC ACGACGTTGT AAAACGACGG CCAGTGAAAT
401 CGGCAATGGG TCGAAATTCA TAGAATTTTG TGTGAGGTGC GTAGCGGCTC
451 TGACAGGGGT GCTGCGCGGA GATCTCTGGT CTCAGGTAGG GCGACAATGG
501 AGAGGTGTTA GTTGCCCCCT GTATCGCTCT CTGCGTGGCG CATTTGGGTCA
551 TCCTGCCCCG ACATATGATA TTCCGCTAGA GGATTACTGA TAGTTTCTGC
601 CTGTCGGGCT TGTCGGGCTT GTCGGGCTTG TCGGGCTTGT CGGGCCTGTC
651 CCTCTTGTC CGCCTGTCT CACTTTTTC CAATCAAAAA ATGGGCGAAG
701 CCCTTCTTGT TCTATAGTTC TTATAGTTCA TACGAAAATT ACACATAATT
751 ATCAATAGCT TATTCGCTTA AAAGGGAGTA ATTGGGCCGC AAAAGGGAGT
801 AATTGGGCCG CAAAAGGGAG TAATTGGGCC GCAAAGGGA GTAATTGGGC
851 CGATATCGGT TGTTTACATG GGGAGGAATC CCCTTAATCA TTTCTCCCCA
901 TGGGAAAGAC AACACAAGTG GCCGCAGACC GGGCCTTCGA CCAGACAAAA
951 ACTGTGCTCC CTGCCCAGGT GCGGAGAGGG GTCTATATGC GCAATCCGCC
1001 CCGCCTGCAG GCGCTCAAGC TCATGCATTT AATGATAGCC ACTGCGGGCG
1051 GCCGCATGGC TGATGATGTG CGCCATGAAA TGCGGCTGGC CGACATTTCG
1101 GCAATCGACG GCATGAAAAA CCATGACCGT GAGAGCCTGA CCCCCTGTT
1151 CGAGGAGCTA GCCGCTGCGG TGTTGACCCA TGATGACCCT GCAAAGATGA
1201 TCGTGACAGT CGGCGGCTTG GTCGATGAGG CGCGAATAGA CTACCGCCAG
1251 GAGGCAAGCG GCGAACTCCT AGTGACGTGG ACCTTCCGGA GTACATTCCG
1301 TCGTATGGCG GCGGAGTCGA ACCACTGGGC CATTCTCGAC CGTCAAACGG
1351 TATTCCATCT CGGTAGTAAG TATTCCGTGC TGCTGTTCCA GCACGTCTCT
1401 AGTCTCGCCA ATCTTGATCG GATGAGCGCG AAAACCTTTA CGGTCCCCGA
1451 GTTGCGGGCG CTCCTTGGAG TGCCCGAGGG AAAGATGGTT CGTTGGAACG
1501 ACGTTAACAG ATTTGCTCTC AAACCTGCAC TGGATGAGAT CAACCATTTA
1551 TCGCGTCTGA CATTGACGGC AAAGCCGACC AAGATTGGCC GTAGCGTGGC
1601 AAGTGTGACT ATAGGCTGGG AAGTGAAAGA CGACCAACC GTCGCCAGGC
1651 GCGAGCTGGC GGGTTCCAAG GTCGGTCGAG ATGCTCGTCG CAGAGGGGCA
1701 GCGGAAACGA TAGCCCCCTC CTTCCAGAA GCGGGCGGGA TCACCTACAG
1751 TCCACGTTGG CTGGAGCTGA AACGCTCTGC TGGCAGCAAC AAGGACAACG
1801 ATCTGATCGC CTCAGACTTC CGGCGTTTCT GTCGGGAGAG AGGCGTGCGT
1851 CTGGACGCTG CAAACATCGA AAAACTGTTT TTAGATTTCT GCGCAAAGGT
1901 AGGGAAGGTT TGAGTTTTGA GGTATTTTAC CGCAATAGTG TTAAATGACT
1951 TTCGTGAAAC GATGTGCAAT ATAGCGGTAA GACTATGAAA TACACGGCTG
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**FIGURE 3A**

2001 GACAGGCTGC AAAAGCAACG GGTGTGGCGA CCGCAACCAT CACTCGGGCG  
2051 CTAAAAAGCG GTAAAAATTC CGGTAAAAAA GATGAATCTG GGGCATGGGT  
2101 TATAGATCCT GCAGAATTGC ACAGAGTGTT TCCTCCCATT TCAAAGAAAT  
2151 ACACCGAAAC ACCTAACACG CAAGTATATG GTAAGCGTGA TGAAACACAT  
2201 GAAATGACCT CAGAAATCAG CGCATTAGAG CGTGAAGTTC GGACTTTACG  
2251 CGATGCTTTA TCTGATGCCA GGGAGGATCG CGACAAATGG CGCGACATGG  
2301 CCGAGCGTCT TTCAATTTCA TCACCGATGA GAGAGGAAGA CCGCCCCCT  
2351 CAAAAACAAA GATGGTGGA GATATTCTGA TCCTGGGCTT CAGGAGCCTT  
2401 GCCTTTACTG GCGGAAAAAC GCGATATTGA GGCACAGGCC CGCACTTTAG  
2451 AGGCGGAAGC CTATAACGAG TACCAAAACA CTAGAAGCCA GATTGAGGAA  
2501 AATAGGGAAC GTGGGATCCT CTAGAGTCGA CCTGCAGGCA TGCAAGCTTG  
2551 GCGTAATCAT GGTCATAGCT GTTTCCTGTG TGAAATTGTT ATCCGCTCAC  
2601 AATTCCACAC AACATACGAG CCGGAAGCAT AAAGTGTAAG GCCTGGGGTG  
2651 CCTAATGAGT GAGCTAACTC ACATTAATTG CGTTGCGCTC ACTGCCCGCT  
2701 TTCCAGTCGG GAAACCTGTC GTGCCAGCTG CATTAATGAA TCGGCCAACG  
2751 CGCGGGGAGA GGCGGTTTGC GTATTGGGCG CTCTTCCGCG CTCGGTCTTG  
2801 CCTTGCTCGT CGGTGATGTA CTTACCAGC TCCGCGAAGT CGCTCTTCTT  
2851 GATGGAGCGC ATGGGGACGT GCTTGGCAAT CACGCGCACC CCCC GGCCGT  
2901 TTTAGCGGCT AAAAAAGTCA TGGCTCTGCC CTCGGGCGGA CCACGCCCAT  
2951 CATGACCTTG CCAAGCTCGT CCTGCTTCTC TTCGATCTTC GCCAGCAGGG  
3001 CGAGGATCGT GGCATCACCG AACCGCGCCG TGCGCGGGTC GTCGGTGAGC  
3051 CAGAGTTTCA GCAGGCCGCC CAGGCGGCC AGGTCGCCAT TGATGCGGGC  
3101 CAGCTCGCGG ACGTGCTCAT AGTCCACGAC GCCCGTGATT TTGTAGCCCT  
3151 GGCCGACGGC CAGCAGGTAG GCCGACAGGC TCATGCCGGC CGCCGCCGCC  
3201 TTTTCTCAA TCGCTCTTCG TTCGCTGGA AGGCAGTACA CCTTGATAGG  
3251 TGGGCTGCCC TTCCTGGTTG GCTTGGTTTC ATCAGCCATC CGCTTGCCCT  
3301 CATCTGTTAC GCCGGCGGTA GCCGGCCAGC CTCGCAGAGC AGGATTCCCCG  
3351 TTGAGCACCG CCAGGTGCGA ATAAGGGACA GTGAAGAAGG AACACCCGCT  
3401 CGCGGGTGGG CTTACTTCAC CTATCCTGCC CGGCTGACGC CGTTGGATAC  
3451 ACCAAGGAAA GTCTACACGA ACCCTTTGGC AAAATCCTGT ATATCGTGCG  
3501 AAAAAGGATG GATATACCGA AAAAATCGCT ATAATGACCC CGAAGCAGGG  
3551 TTATGCAGCG GAAAAGCTTC CTCGCTCACT GACTCGCTGC GCTCGGTCGT  
3601 TCGGCTGCGG CGAGCGGTAT CAGCTCACTC AAAGGCGGTA ATACGGTTAT  
3651 CCACAGAATC AGGGGATAAC GCAGGAAAGA ACATGTGAGC AAAAGGCCAG  
3701 CAAAAGGCCA GGAACCGTAA AAAGGCCGCG TTGCTGGCGT TTTTCCATAG  
3751 GCTCCGCCCC CCTGACGAGC ATCACAAAAA TCGACGCTCA AGTCAGAGGT  
3801 GGCGAAACCC GACAGGACTA TAAAGATACC AGGCGTTTCC CCCTGGAAGC  
3851 TCCCTCGTGC GCTCTCCTGT TCCGACCCTG CCGCTTACCG GATACCTGTC  
3901 CGCCTTTCTC CCTTCGGGAA GCGTGGCGCT TTCTCATAGC TCACGCTGTA  
3951 GGTATCTCAG TTCGGTGTAG GTCGTTGCT CCAAGCTGGG CTGTGTGCAC  
4001 GAACCCCCCG TTCAGCCCGA CCGCTGCGCC TTATCCGGTA ACTATCGTCT  
4051 TGAGTCCAAC CCGGTAAGAC ACGACTTATC GCCACTGGCA GCAGCCACTG

**FIGURE 3B**

4101 GTAACAGGAT TAGCAGAGCG AGGTATGTAG GCGGTGCTAC AGAGTTCTTG  
 4151 AAGTGGTGGC CTAACCTACGG CTACACTAGA AGGACAGTAT TTGGTATCTG  
 4201 CGCTCTGCTG AAGCCAGTTA CCTTCGGAAA AAGAGTTGGT AGCTCTTGAT  
 4251 CCGGCAAAACA AACCACCGCT GGTAGCGGTG GTTTTTTTGT TTGCAAGCAG  
 4301 CAGATTACGC GCAGAAAAAA AGGATCTCAA GAAGATCCTT TGATCTTTTC  
 4351 TACGGGGTCT GACGCTCAGT GGAACGAAAA CTCACGTAA GGGATTTTGG  
 4401 TCATGAGATT ATCAAAAAGG ATCTTCACCT AGATCCTTTT GGGGGGGGGG  
 4451 GGCGCTGAGG TCTGCCTCGT GAAGAAGGTG TTGCTGACTC ATACCAGGCC  
 4501 TGAATCGCCC CATCATCCAG CCAGAAAGTG AGGGAGCCAC GGTTGATGAG  
 4551 AGCTTTGTTG TAGGTGGACC AGTTGGTGAT TTTGAACCTT TGCTTTGCCA  
 4601 CGGAACGGTC TCGCTTGTCG GGAAGATGCG TGATCTGATC CTTCAACTCA  
 4651 GCAAAAGTTC GATTTATTCA ACAAAGCCGC CGTCCCGTCA AGTCAGCGTA  
 4701 ATGCTCTGCC AGTGTTACAA CCAATTAACC AATTCTGATT AGAAAACTC  
 4751 ATCGAGCATC AAATGAAACT GCAATTTATT CATATCAGGA TTATCAATAC  
 4801 CATATTTTTG AAAAAGCCGT TTCTGTAATG AAGGAGAAAA CTCACCGAGG  
 4851 CAGTTCCTATA GGATGGCAAG ATCCTGGTAT CGGTCTGCGA TTCCGACTCG  
 4901 TCCAACATCA ATACAACCTA TTAATTTCCC CTCGTCAAAA ATAAGGTTAT  
 4951 CAAGTGAGAA ATCACCATGA GTGACGACTG AATCCGGTGA GAATGGCAAA  
 5001 AGCTTATGCA TTTCTTTCCA GACTTGTTCA ACAGGCCAGC CATTACGCTC  
 5051 GTCATCAAAA TCACTCGCAT CAACCAAACC GTTATTCATT CGTGATTGCG  
 5101 CCTGAGCGAG ACGAAATACG CGATCGCTGT TAAAAGGACA ATTACAAACA  
 5151 GGAATCGAAT GCAACCGGCG CAGGAACACT GCCAGCGCAT CAACAATATT  
 5201 TTCACCTGAA TCAGGATATT CTTCTAATAC CTGGAATGCT GTTTTCCCGG  
 5251 GGATCGCAGT GGTGAGTAAC CATGCATCAT CAGGAGTACG GATAAAATGC  
 5301 TTGATGGTCG GAAGAGGCAT AAATTCCGTC AGCCAGTTTA GTCTGACCAT  
 5351 CTCATCTGTA ACATCATTGG CAACGCTACC TTTGCCATGT TTCAGAAACA  
 5401 ACTCTGGCGC ATCGGGCTTC CCATACAATC GATAGATTGT CGCACCTGAT  
 5451 TGCCCGACAT TATCGCGAGC CCATTTATAC CCATATAAAT CAGCATCCAT  
 5501 GTTGGAATTT AATCGCGGCC TCGAGCAAGA CGTTTCCCGT TGAATATGGC  
 5551 TCATAACACC CCTTGTATTA CTGTTTATGT AAGCAGACAG TTTTATTGTT  
 5601 CATGATGATA TATTTTTATC TTGTGCAATG TAACATCAGA GATTTTGAGA  
 5651 CACAACGTGG CTTTCCCCC CCCCCATTA TTGAAGCATT TATCAGGGTT  
 5701 ATTGTCTCAT GAGCGGATAC ATATTTGAAT GTATTTAGAA AAATAAACAA  
 5751 ATAGGGGTTC CGCGCACATT TCCCCGAAAA GTGCCACCTG ACGTCTAAGA  
 5801 AACCATTATT ATCATGACAT TAACCTATAA AAATAGGCGT ATCACGAGGC  
 5851 CCTTTCGTC

**FIGURE 3C**

SEQ ID NO:4

1 GGCAATGGGT CGAAATTCAT AGAATTTTGT GTGAGGTGCG TAGCGGCTCT  
51 GACAGGGGTG CTGCGCGGAG ATCTCTGGTC TCAGGTAGGG CGACAATGGA  
101 GAGGTGTTAG TTGCCCCCTG TATCGCTCTC TGCCTGGCGC ATTGGGTCAT  
151 CCTGCCCCGA CATATGATAT TCCGCTAGAG GATTACTGAT AGTTTCTGCC  
201 TGTCTGGGCTT GTCGGGCTTG TCGGGCTTGT CGGGCTTGTC GGGCCTGTCC  
251 CTCTTGTCCC GCCTGTCCTC ACTTTTTCAC AATCAAAAAA TGGGCGAAGC  
301 CCTTCTTGTT CTATAGTTCT TATAGTTCAT ACGAAAATTA CACATAATTA  
351 TCAATAGCTT ATTCGCTTAA AAGGGAGTAA TTGGGCCGCA AAAGGGAGTA  
401 ATTGGGCCGC AAAAGGGAGT AATTGGGCCG CAAAAGGGAG TAATTGGGCC  
451 GATATCGGTT GTTTACATGG GGAGGAATCC CCTTAATCAT TTCTCCCCAT  
501 GGGAAAGACA ACACAAGTGG CCGCAGACCG GGCTTCGAC CAGACAAAAA  
551 CTGTGCTCCC TGCCGAGGTG GCGAGAGGGG TCTATATGCG CAATCCGCCC  
601 CGCTGCAGG CGCTCAAGCT CATGCATTTA ATGATAGCCA CTGCGGGCGG  
651 CCGCATGGCT GATGATGTGC GCCATGAAAT GCGGCTGGCC GACATTGCGG  
701 CAATCGACGG CATGAAAAAC CATGACCGTG AGAGCCTGAC CCCGCTGTTC  
751 GAGGAGCTAG CCGCTGCGGT GTTGACCCAT GATGACCCCTG CAAAGATGAT  
801 CGTGACAGTC GGCGGCTTGG TCGATGAGGC GCGAATAGAC TACCGCCAGG  
851 AGGCAAGCGG CGAACTCCTA GTGACGTGGA CCTTCCGGAG TACATTCCGT  
901 CGTATGGCGG CGGAGTCGAA CCACTGGGCC ATTCTCGACC GTCAAACGGT  
951 ATTCCATCTC GGTAGTAAGT ATTCCGTGCT GCTGTTCCAG CACGTCTCTA  
1001 GTCTCGCCAA TCTTGATCGG ATGAGCGCGA AAACCTTTAC GGTCCCCGAG  
1051 TTGCGGGCGC TCCTTGAGT GCGCGAGGGA AAGATGGTTC GTTGGAACGA  
1101 CGTTAACAGA TTTGCTCTCA AACCTGCACT GGATGAGATC AACCATTTAT  
1151 CCGTCTGAC ATTGACGGCA AAGCCGACCA AGATTGGCCG TAGCGTGGA  
1201 AGTGTGACTA TAGGCTGGGA AGTGAAAGAC GACCCAACCG TCGCCAGGCG  
1251 CGAGCTGGCG GGTTCCAAGG TCGGTCGAGA TGCTCGTCGC AGAGGGGCAG  
1301 CGGAAACGAT AGCCCCCTCC TTCCCAGAAG CGGGCGGGAT CACCTACAGT  
1351 CCACGTTGGC TGGAGCTGAA ACGCTCTGCT GGCAGCAACA AGGACAACGA  
1401 TCTGATCGCC TCAGACTTCC GCGCTTTCTG TCGGGAGAGA GCGGTGCGTC  
1451 TGGACGCTGC AAACATCGAA AAAGTGTGTT TAGATTCTCTG CGCAAAGGTA  
1501 GGGAAAGGTTT GAGTTTTGAG GTATTTTACC GCAATAGTGT TAAATGACTT  
1551 TCGTGAAACG ATGTGCAATA TAGCGGTAAG ACTATGAAAT ACACGGCTGG  
1601 ACAGGCTGCA AAAGCAACGG GTGTGGCGAC CGCAACCATC ACTCGGGCGC  
1651 TAAAAAGCGG TAAATTTTCC GGTAAAAAAG ATGAATCTGG GGCATGGGTT  
1701 ATAGATCCTG CAGAATTGCA CAGAGTGTTT CCTCCCATT CAAAGAAATA  
1751 CACCGAAACA CCTAACACGC AAGTATATGG TAAGCGTGAT GAAACACATG  
1801 AAATGACCTC AGAAATCAGC GCATTAGAGC GTGAAGTTCG GACTTTACGC  
1851 GATGCTTTAT CTGATGCCAG GGAGGATCGC GACAAATGGC GCGACATGGC  
1901 CGAGCGTCTT TCAATTTTAT CACCGATGAG AGAGGAAGAC CGCCCCCTC  
1951 AAAAACAAAG ATGGTGGAAG ATATTCTGAT CCTGGGCTTC AGGAGCCTTG

FIGURE 4A

